REMARKS

Applicants respectfully request reconsideration of the present application in view of the foregoing amendments.

Applicants filed a Sequence Listing on January 30, 2003, after they incorporated the actual DNA and amino acid sequences of GenBank Accession No. Y08612 into the specification and claims in their paper of October 10, 2002. The present Amendment now incorporates the specific sequence identifiers of those sequences in the specification and claims. Thus, page 6 of the specification, which describes the exact DNA and amino acid sequences of Y08612, now includes references "(SEQ ID NO. 1)" for the DNA sequence and "(SEQ ID NO. 2)" for the amino acid sequence.

Similarly, claims 3, 23, and 24, have been amended to recite "the protein consisting of the amino acid sequence of SEQ ID NO. 2."

The amendment is fully supported by the specification and Applicants have not introduced any new matter into the claims.

IV. Conclusion

Applicants believe that the claimed invention is now in condition for allowance. Favorable reconsideration of the application as amended is respectfully requested. The Examiner is invited to contact the undersigned by telephone if it is felt that a telephone interview would advance the prosecution of the present application.

Respectfully submitted,

Date Feb 5, 2003

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MARKED-UP VERSION OF THE SPECIFICATION

At page 6, please add the following paragraphs before the "Brief Description of the Drawings":

The amino acid sequence (SEQ ID NO. 2) of accession number Y08612 is:

MAAAEGPVGDGELWQTWLPNHVVFLRLREGLKNQSPTEAEKPASSSLPSSPPPQLLTRNVVFGLGGELFLWDGED SSFLVVRLRGPSGGGEEPALSQYQRLLCINPPLFEIYQVLLSPTQHHVALIGIKGLMVLELPKRWGKNSEFEGGK STVNCSTTPVAERFFTSSTSLTLKHAAWYPSEILDPHVVLLTSDNVIRIYSLREPQTPTNVIILSEAEEESLVLN KGRAYTASLGETAVAFDFGPLDAVPKTLFGQNGKDEVVAYPLYILYENGETFLTYISLLHSPGNIWKAVGSIAHA SAAEDNYGYDACAVLCLPCVPNILVIATESGMLYHCVVLEGEEEDDHTSEKSWDSRIDLIPSLYVFECVELELAL KLASGEDDPFDSDFSCPVKLHRDPKCPSRYHCTHEAGVHSVGLTWIHKLHKFLGSDEEDKDSLQELSTEQKCFVE HILCTRPLPCRQPAPIRGFWIVPDILGPTMICITSTYECLIWPLLSTVHPASPPLLCTREDVEVAESSLRVLAET PDSFEKHIRSILQRSVANPAFLKASEKDIAPPPEECLQLLSRATQVFREQYILKQDLAKEEIQRRVKLLCDQKKK QLEDLSYCREERKSLREMAERLADKYEEAKEKQEDIMNRMKKLLHSFHSELPVLSDSERDMKKELQLIPDQLRHL GNAIKQVTMKKDYQQQKMEKVLSLPKPTIILSAYQRKCIQSILKEEGEHIREMVKQINDIRNHVNF.

The nucleic acid sequence (SEQ ID NO. 1) encoding the protein of accession number Y08612 is:

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1 gataaaccca caagacacaa aacatacctt tcgagcagtt gggccaagat ggcggccgcc
  61 gagggaccgg tgggcgacgg cgagctgtgg cagacctggc ttcctaacca cgtcgtgttc
 121 ttgcqqctcc qqqaqqqact qaaaaaccaq agtccaaccq aagctqagaa accagcttct
 181 togtogttgc cttcgtcgcc gccgccgcag ttgctgacga gaaacgtggt ctttggcctc
 241 ggcggagagc ttttcctgtg ggacggagaa gacagctcct tcttagtcgt tcgccttcgg
 301 ggccccagcg gcggcggcga agagcccgcc ctgtcccagt accagagatt gctttgcata
 421 cttataggaa taaaaggact tatggtatta gaattaccta aaagatgggg gaagaattct
 481 gaatttgaag gtggaaaatc aacagtgaat tgtagtacca ctccagttgc ggagagattt
 541 ttcaccagtt ccacctctct gactctaaag catgctgcat ggtatccaag tgaaatcctg
 601 gatccccacg tagtgctgtt aacatcagac aacgtaatca gaatttactc tctacgtgag
 661 ccgcagacac ccactaacgt gataatactt tcagaagccg aagaggaaag tctagtactc
 721 aataaaggaa gggcgtatac cgcatctcta ggagagacag cagttgcatt tgactttggg
 781 ccattggacg cagtcccaaa gactctattt ggacaaaacg gcaaagatga agtagtggca
 841 tacccactgt acatcttata tgaaaatgga gagactttcc tgacatacat cagtctgtta
 901 cacagooctg gaaatatttg gaaagotgtt gggtccattg cocatgoatc tgcggctgaa
 961 gataactatg gttatgatgc gtgtgctgta ctctgcttac cctgtgtccc caatatctta
1021 gtgatcgcta ctgaatcagg aatgctgtat cactgtgtcg tgctagaagg ggaagaagaa
1081 gatgaccaca cgtcagaaaa gtcctgggat tccaggattg acctcattcc ttctctgtat
1141 gtgtttgaat gtgttgagtt ggagcttgct ttgaaactgg catctggaga ggatgaccct
1201 tttgattctg acttttcttg tccagtcaaa cttcatagag atcccaagtg tccttcaaga
1261 tatcactgta ctcatgaagc tggtgtacat agtgttgggc taacttggat tcataaactt
1321 cacaaatttc ttggatcaga tgaagaagat aaggatagtt tacaggaact ctctacagaa
1381 cagaaatgct ttgttgaaca catcetttgt acgaggccat tgccctgcag gcagccagct
1441 ccaattcgag gattttggat tgtacctgac attctgggac ccacgatgat ctgcatcacc
1501 agtacctatg aatgcctcat atggccgtta ttaagtacag tccatccagc gtctcctccc
1561 ctgctttgta ctcgagaaga tgttgaagtg gcagagtctt ccctccgtgt tctggctgaa
1621 accccagatt cctttgaaaa gcatattaga agcattttgc aacgtagtgt tgccaatcca
1681 gcatttttga aagcttctga aaaggacata gcccctcctc ctgaagaatg ccttcagctc
1741 ctcagcagag ccacccaggt gttcagagag cagtacattc tcaaacagga cttggcaaag
1801 gaggagattc agcggagggt caaattatta tgtgaccaaa aaaagaaaca actagaagat
1861 ctcagttatt gtcgagaaga gaggaaaagt ctgcgggaaa tggctgagcg tttagctgac
1921 aaatatqaqq aaqctaaaga aaaacaaqag gatatcatga acaggatgaa aaaactactt
1981 cacagttttc actctgagct cccagttctc tctgatagtg agcgagacat gaagaaagaa
2041 ttacagctga tacctgatca acttcgacat ttgggcaatg ccatcaaaca ggttactatg
2101 aaaaaggatt atcaacagca aaagatggag aaggtgttga gtcttccaaa acccaccatt
2161 attctcagtg cctaccagcg aaagtgcatt cagtccatcc tgaaagagga gggtgaacat
2221 ataagggaaa tggtgaagca aatcaatgat atccgcaatc atgtaaactt ctgacaccac
2281 caggagetga eteacacetg aactgaacac cattgaagge ttaaacccat attgtaaaac
2341 aggtagaatt atctaattta taaaaaggtg ttttgatg.
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MARKED-UP VERSION OF THE CLAIMS

- 13. (Twice amended) A method for identifying a cancer cell comprising:
 - (a) providing a tissue biopsy sample; and
- (b) determining the level of expression in said sample of the protein consisting of the amino acid sequence of SEQ ID NO. 2:

MAAAEGPVGDGELWQTWLPNHVVFLRLREGLKNQSPTEAEKPASSSLPSSPPPQLLTRNVVFGLGGELFLWDGE DSSFLVVRLRGPSGGEEPALSQYQRLLCINPPLFEIYQVLLSPTQHHVALIGIKGLMVLELPKRWGKNSEFEG GKSTVNCSTTPVAERFFTSSTSLTLKHAAWYPSEILDPHVVLLTSDNVIRIYSLREPQTPTNVIILSEAEEESL VLNKGRAYTASLGETAVAFDFGPLDAVPKTLFGQNGKDEVVAYPLYILYENGETFLTYISLLHSPGNIWKAVGS IAHASAAEDNYGYDACAVLCLPCVPNILVIATESGMLYHCVVLEGEEEDDHTSEKSWDSRIDLIPSLYVFECVE LELALKLASGEDDPFDSDFSCPVKLHRDPKCPSRYHCTHEAGVHSVGLTWIHKLHKFLGSDEEDKDSLQELSTE QKCFVEHILCTRPLPCRQPAPIRGFWIVPDILGPTMICITSTYECLIWPLLSTVHPASPPLLCTREDVEVAESS LRVLAETPDSFEKHIRSILQRSVANPAFLKASEKDIAPPPEECLQLLSRATQVFREQYILKQDLAKEEIQRRVK LLCDQKKKQLEDLSYCREERKSLREMAERLADKYEEAKEKQEDIMNRMKKLLHSFHSELPVLSDSERDMKKELQ LIPDQLRHLGNAIKQVTMKKDYQQQKMEKVLSLPKPTIILSAYQRKCIQSILKEEGEHIREMVKQINDIRNHVN F,

wherein a sample comprising said protein at a level of expression that is greater than non-cancer cells indicates that said sample comprises a cancer cell.

23. (Twice amended) A diagnostic kit comprising a protein binding molecule, wherein the protein binding molecule binds to the protein consisting of the amino acid sequence of SEQ ID NO. 2:

MAAAEGPVGDGELWQTWLPNHVVFLRLREGLKNQSPTEAEKPASSSLPSSPPPQLLTRNVVFGLGGELFLWDGED SSFLVVRLRGPSGGGEEPALSQYQRLLCINPPLFEIYQVLLSPTQHHVALIGIKGLMVLELPKRWGKNSEFEGGK STVNCSTTPVAERFFTSSTSLTLKHAAWYPSEILDPHVVLLTSDNVIRIYSLREPQTPTNVIILSEAEEESLVLN KGRAYTASLGETAVAFDFGPLDAVPKTLFGQNGKDEVVAYPLYILYENGETFLTYISLLHSPGNIWKAVGSIAHA SAAEDNYGYDACAVLCLPCVPNILVIATESGMLYHCVVLEGEEEDDHTSEKSWDSRIDLIPSLYVFECVELELAL KLASGEDDPFDSDFSCPVKLHRDPKCPSRYHCTHEAGVHSVGLTWIHKLHKFLGSDEEDKDSLQELSTEQKCFVE HILCTRPLPCRQPAPIRGFWIVPDILGPTMICITSTYECLIWPLLSTVHPASPPLLCTREDVEVAESSLRVLAET PDSFEKHIRSILQRSVANPAFLKASEKDIAPPPEECLQLLSRATQVFREQYILKQDLAKEEIQRRVKLLCDQKKK QLEDLSYCREERKSLREMAERLADKYEEAKEKQEDIMNRMKKLLHSFHSELPVLSDSERDMKKELQLIPDQLRHL GNAIKQVTMKKDYQQQKMEKVLSLPKPTIILSAYQRKCIQSILKEEGEHIREMVKQINDIRNHVNF.

24. (Twice amended) A diagnostic kit comprising a nucleic acid, wherein the nucleic acid anneals specifically to a nucleic acid transcript that encodes the protein consisting of the amino acid sequence of SEQ ID NO. 2:

MAAAEGPVGDGELWQTWLPNHVVFLRLREGLKNQSPTEAEKPASSSLPSSPPPQLLTRNVVFGLGGELFLWDGED SSFLVVRLRGPSGGGEEPALSQYQRLLCINPPLFEIYQVLLSPTQHHVALIGIKGLMVLELPKRWGKNSEFEGGK STVNCSTTPVAERFFTSSTSLTLKHAAWYPSEILDPHVVLLTSDNVIRIYSLREPQTPTNVIILSEAEEESLVLN KGRAYTASLGETAVAFDFGPLDAVPKTLFGQNGKDEVVAYPLYILYENGETFLTYISLLHSPGNIWKAVGSIAHA SAAEDNYGYDACAVLCLPCVPNILVIATESGMLYHCVVLEGEEEDDHTSEKSWDSRIDLIPSLYVFECVELELAL KLASGEDDPFDSDFSCPVKLHRDPKCPSRYHCTHEAGVHSVGLTWIHKLHKFLGSDEEDKDSLQELSTEQKCFVE HILCTRPLPCRQPAPIRGFWIVPDILGPTMICITSTYECLIWPLLSTVHPASPPLLCTREDVEVAESSLRVLAET PDSFEKHIRSILQRSVANPAFLKASEKDIAPPPEECLQLLSRATQVFREQYILKQDLAKEEIQRRVKLLCDQKKK QLEDLSYCREERKSLREMAERLADKYEEAKEKQEDIMNRMKKLLHSFHSELPVLSDSERDMKKELQLIPDQLRHL GNAIKQVTMKKDYQQQKMEKVLSLPKPTIILSAYQRKCIQSILKEEGEHIREMVKQINDIRNHVNF.